

RAW SEQUENCE LISTING

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Application Serial Number: 10/S16,088B
Source: PCT
Date Processed by STIC: 4-28-06

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DATE: 04/28/2006

PATENT APPLICATION: US/10/516,088B

TIME: 09:43:05

Input Set : A:\corrected P70284US0.APP

Output Set: N:\CRF4\04282006\J516088B.raw

3 <110> APPLICANT: BRANCIA, FRANCESCO LORENZO
 5 <120> TITLE OF INVENTION: SELECTIVE BINDING AND ANALYSIS OF MACROMOLECULES
 7 <130> FILE REFERENCE: 12687/P70284US0
 9 <140> CURRENT APPLICATION NUMBER: US/10/516,088B
 10 <141> CURRENT FILING DATE: 2004-11-30
 12 <150> PRIOR APPLICATION NUMBER: GB0212522.7
 13 <151> PRIOR FILING DATE: 2002-05-31
 15 <160> NUMBER OF SEQ ID NOS: 4
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 7
 21 <212> TYPE: PRT
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Synthetic peptide made by New England Peptide,
 Inc. Gardner, MA;
 26 Lysine terminal standard peptide AFLDASK
 28 <400> SEQUENCE: 1
 30 Ala Phe Leu Asp Ala Ser Lys
 31 1 5
 33 <210> SEQ ID NO: 2
 34 <211> LENGTH: 7
 35 <212> TYPE: PRT
 36 <213> ORGANISM: Artificial Sequence
 38 <220> FEATURE:
 39 <223> OTHER INFORMATION: Synthetic peptide made by New England Peptide,
 Inc. Gardner, MA;
 40 Homoarginine terminal standard peptide made by guanidination of
 41 the terminal Lys and identified in the specification as AFLDASK*
 43 <400> SEQUENCE: 2
 45 Ala Phe Leu Asp Ala Ser Arg
 46 1 5
 48 <210> SEQ ID NO: 3
 49 <211> LENGTH: 7
 50 <212> TYPE: PRT
 51 <213> ORGANISM: Artificial Sequence
 53 <220> FEATURE:
 54 <223> OTHER INFORMATION: Synthetic peptide made by New England Peptide,
 Inc. Gardner, MA;
 55 Arginine terminal standard peptide AFLDASR
 57 <400> SEQUENCE: 3
 59 Ala Phe Leu Asp Ala Ser Arg
 60 1 5
 62 <210> SEQ ID NO: 4

63 <211> LENGTH: 9
64 <212> TYPE: PRT

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65 <213> ORGANISM: Artificial Sequence

67 <220> FEATURE:

68 <222> LOCATION: 163..171

69 <223> OTHER INFORMATION: Interleukin 1 beta, Homo Sapiens; Synthetic peptide made by

70 Sigma, Inc.

72 <400> SEQUENCE: 4

74 Val Gln Gly Glu Glu Ser Asn Asp Lys

75 1 5

VERIFICATION SUMMARY

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